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1: NM_017831. Homo sapiens hypo...[gi:8923428]

Related Sequences, Protein, Taxonomy, LinkOut

LOCUS NM_017831 1916 bp mRNA linear PRI 10-DEC-2001
 DEFINITION Homo sapiens hypothetical protein FLJ20456 (FLJ20456), mRNA.
 ACCESSION NM_017831
 VERSION NM_017831.1 GI:8923428
 KEYWORDS .
 SOURCE human.

2/15/00

ORGANISM Homo sapiens
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
 Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

REFERENCE 1 (sites)

AUTHORS Watanabe,K., Kumagai,A., Itakura,S., Yamazaki,M., Tashiro,H.,
 Ota,T., Suzuki,Y., Obayashi,M., Nishi,T., Shibahara,T., Tanaka,T.,
 Nakamura,Y., Isogai,T. and Sugano,S.

TITLE NEDO human cDNA sequencing project

JOURNAL Unpublished (2000)

COMMENT PREDICTED REFSEQ: The mRNA record is supported by experimental
 evidence; however, the coding sequence is predicted. The reference
 sequence was derived from AK000463.1.

FEATURES Location/Qualifiers

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1: AK000463. Homo sapiens cDNA...[gi:7020568]

Related Sequences, Protein, Taxonomy, LinkOut

LOCUS AK000463 1916 bp mRNA linear PRI 22-FEB-2000
 DEFINITION Homo sapiens cDNA FLJ20456 fis, clone KAT05827.
 ACCESSION AK000463
 VERSION AK000463.1 GI:7020568
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 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
 REFERENCE 1 (sites)
 AUTHORS Watanabe,K., Kumagai,A., Itakura,S., Yamazaki,M., Tashiro,H., Ota,T., Suzuki,Y., Obayashi,M., Nishi,T., Shibahara,T., Tanaka,T., Nakamura,Y., Isogai,T. and Sugano,S.
 TITLE NEDO human cDNA sequencing project
 JOURNAL Unpublished (2000)
 REFERENCE 2 (bases 1 to 1916)
 AUTHORS Sugano,S., Suzuki,Y., Ota,T., Obayashi,M., Nishi,T., Isogai,T., Shibahara,T., Tanaka,T. and Nakamura,Y.
 TITLE Direct Submission
 JOURNAL Submitted (15-FEB-2000) Sumio Sugano, Institute of Medical Science, University of Tokyo, Deptment of Virology; Shirokane-dai, 4-6-1, Minato-ku, Tokyo 108-8639, Japan (E-mail:cdna1@ims.u-tokyo.ac.jp, Tel:81-3-5449-5286, Fax:81-3-5449-5416)
 COMMENT NEDO human cDNA sequencing project supported by Ministry of International Trade and Industry of Japan; cDNA full insert sequencing: Research Association for Biotechnology; cDNA library construction, 5'- & 3'-end one pass sequencing: Department of Virology and Human Genome Center, Institute of Medical Science, University of Tokyo (partly supported by Science and Technology Agency).
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1: XM_008732. Homo sapiens hypo...[gi:18588856]

Related Sequences. Protein, Taxonomy, LinkOut

LOCUS XM_008732 1916 bp mRNA linear PRI 07-FEB-2002
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 VERSION XM_008732.4 GI:18588856
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 Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
 REFERENCE 1 (bases 1 to 1916)
 AUTHORS NCBI Annotation Project.
 TITLE Direct Submission
 JOURNAL Submitted (06-FEB-2002) National Center for Biotechnology
 Information, NIH, Bethesda, MD 20894, USA
 COMMENT GENOME ANNOTATION REFSEQ: This model reference sequence was
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 analysis using gene prediction method: BLAST. -Also see:-
 Documentation of NCBI's Annotation Process- Evidence Viewer -
 alignments supporting this model.
 On Feb 7, 2002 this sequence version replaced gi:17456245.
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